

ABSTRACT OF THE DISCLOSURE

A method for determining whether a compound inhibits or disrupts an interaction between a first polypeptide comprising a transcriptional adaptor motif (TRAM) and a second polypeptide comprising a TRAM-interaction motif. The first polypeptide and/or second polypeptide may be Mdm-2, p53, TBP, E2F, YY1, CBP/p300 or TF11B, or a viral polypeptide such as a human papillomavirus (HPV) E6 polypeptide from HPV strain (16) or (18).

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